

## Patent Application US/07/996,771

## SEQUENCE LISTING

ENTERED

## (1) GENERAL INFORMATION:

(i) APPLICANT: R. Bruce Wallace

(ii) TITLE OF INVENTION: Ligation Amplification of  
Nucleic Acid Sequences

(iii) NUMBER OF SEQUENCES: 15

## (iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: City of Hope

(B) STREET: 1500 East Duarte Road

(C) CITY: Duarte

(D) STATE: California

(E) COUNTRY: United States of America

(F) ZIP: 91010-0269

## (v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: 3M Double Density 5 1/4" diskette

(B) COMPUTER: Wang PC

(C) OPERATING SYSTEM: MS-DOS (R) Version 3.30

(D) SOFTWARE: Microsoft (R)

## (vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: US/07/996,771

(B) FILING DATE: 24 DEC 1992

(C) CLASSIFICATION: 435

## (vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: US/07/870,221

(B) FILING DATE: 20 April 1992

(C) CLASSIFICATION: Unknown

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53  
54 (D) APPLICATION NUMBER: US/07/178,377  
55  
56 (B) FILING DATE: 06 April 1988  
57  
58 (C) CLASSIFICATION: Unknown  
59  
60 (viii) ATTORNEY/AGENT INFORMATION:  
61  
62 (A) NAME: Irons, Edward S.  
63  
64 (B) REGISTRATION NUMBER: 16,541  
65  
66 (C) REFERENCE/DOCKET NUMBER: None  
67  
68 (ix) TELECOMMUNICATION INFORMATION:  
69  
70 (A) TELEPHONE: (202) 785-6938  
71  
72 (B) TELEFAX: (202) 785-5351  
73  
74 (C) TELEX: 440087 LM WSH  
75  
76  
77 (2) INFORMATION FOR SEQ ID NO: 1:  
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79 (i) SEQUENCE CHARACTERISTICS:  
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81 (A) LENGTH: 14  
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83 (B) TYPE: nucleic acid  
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85 (C) STRANDEDNESS: Single Stranded  
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87 (D) TOPOLOGY: Linear  
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89 (ii) MOLECULE TYPE: Not Applicable  
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91 (iii) HYPOTHETICAL: Not Applicable  
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93 (iv) ANTI-SENSE: Not Applicable  
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95 (v) FRAGMENT TYPE: Not Applicable  
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97 (vi) ORIGINAL SOURCE: Synthetically Prepared  
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99 (vii) IMMEDIATE SOURCE: Synthetically Prepared  
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101 (viii) POSITION IN GENOME: None  
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103 (ix) FEATURE: None  
104

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105 (x) PUBLICATION INFORMATION: None  
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107 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:  
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109 CGGCAGACTT CTCC 14  
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113 (2) INFORMATION FOR SEQ ID NO: 2:  
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115 (i) SEQUENCE CHARACTERISTICS:  
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117 (A) LENGTH: 8  
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119 (B) TYPE: nucleic acid  
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121 (C) STRANDEDNESS: Single Stranded  
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123 (D) TOPOLOGY: Linear  
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125 (ii) MOLECULE TYPE: Not Applicable  
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137 (viii) POSITION IN GENOME: None  
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139 (ix) FEATURE: None  
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141 (x) PUBLICATION INFORMATION: None  
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145 TCAGGAGT 8  
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149 (2) INFORMATION FOR SEQ ID NO:3:  
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151 (i) SEQUENCE CHARACTERISTICS:  
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153 (A) LENGTH: 6  
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155 (B) TYPE: nucleic acid  
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157 (C) STRANDEDNESS: Single Stranded  
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159 (D) TOPOLOGY: Linear  
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161 (ii) MOLECULE TYPE: Not Applicable  
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185 (i) SEQUENCE CHARACTERISTICS:  
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187 (A) LENGTH: 8  
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189 (B) TYPE: nucleic acid  
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191 (C) STRANDEDNESS: Single Stranded  
192  
193 (D) TOPOLOGY: Linear  
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195 (ii) MOLECULE TYPE: Not Applicable  
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197 (iii) HYPOTHETICAL: Not Applicable  
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199 (iv) ANTI-SENSE: Not Applicable  
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201 (v) FRAGMENT TYPE: Not Applicable  
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203 (vi) ORIGINAL SOURCE: Synthetically Prepared  
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205 (vii) IMMEDIATE SOURCE: Synthetically Prepared  
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207 (viii) POSITION IN GENOME: None  
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209 (ix) FEATURE: None  
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211 (x) PUBLICATION INFORMATION: None  
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226 (A) LENGTH: 6  
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228 (B) TYPE: nucleic acid  
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230 (C) STRANDEDNESS: Single Stranded  
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232 (D) TOPOLOGY: Linear  
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234 (ii) MOLECULE TYPE: Not Applicable  
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236 (iii) HYPOTHETICAL: Not Applicable  
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238 (iv) ANTI-SENSE: Not Applicable  
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248 (ix) FEATURE: None  
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250 (x) PUBLICATION INFORMATION: None  
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258 (2) INFORMATION FOR SEQ ID NO:6:  
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260 (i) SEQUENCE CHARACTERISTICS:

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261  
262 (A) LENGTH: 14  
263  
264 (B) TYPE: nucleic acid  
265  
266 (C) STRANDEDNESS: Single Stranded  
267  
268 (D) TOPOLOGY: Linear  
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270 (ii) MOLECULE TYPE: Not Applicable  
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272 (iii) HYPOTHETICAL: Not Applicable  
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274 (iv) ANTI-SENSE: Not Applicable  
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276 (v) FRAGMENT TYPE: Not Applicable  
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278 (vi) ORIGINAL SOURCE: Synthetically Prepared  
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280 (vii) IMMEDIATE SOURCE: Synthetically Prepared  
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298 (A) LENGTH: 14  
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304 (D) TOPOLOGY: Linear  
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306 (ii) MOLECULE TYPE: Not Applicable  
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308 (iii) HYPOTHETICAL: Not Applicable  
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310 (iv) ANTI-SENSE: Not Applicabl  
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312 (v) FRAGMENT TYPE: Not Applicable

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313  
314 (vi) ORIGINAL SOURCE: Synthetically Prepared  
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316 (vii) IMMEDIATE SOURCE: Synthetically Prepared  
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318 (viii) POSITION IN GENOME: None  
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322 (x) PUBLICATION INFORMATION: None  
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330 (2) INFORMATION FOR SEQ ID NO:8:  
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334 (A) LENGTH: 8  
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336 (B) TYPE: nucleic acid  
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338 (C) STRANDEDNESS: Single Stranded  
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340 (D) TOPOLOGY: Linear  
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342 (ii) MOLECULE TYPE: Not Applicable  
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344 (iii) HYPOTHETICAL: Not Applicable  
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346 (iv) ANTI-SENSE: Not Applicable  
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350 (vi) ORIGINAL SOURCE: Synthetically Prepared  
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354 (viii) POSITION IN GENOME: None  
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356 (ix) FEATURE: None  
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358 (x) PUBLICATION INFORMATION: None  
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360 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:  
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362 GGAGAAGT 8  
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368 (i) SEQUENCE CHARACTERISTICS:  
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370 (A) LENGTH: 6  
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372 (B) TYPE: nucleic acid  
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374 (C) STRANDEDNESS: Single Stranded  
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376 (D) TOPOLOGY: Linear  
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378 (ii) MOLECULE TYPE: Not Applicable  
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380 (iii) HYPOTHETICAL: Not Applicable  
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382 (iv) ANTI-SENSE: Not Applicable  
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384 (v) FRAGMENT TYPE: Not Applicable  
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386 (vi) ORIGINAL SOURCE: Synthetically Prepared  
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388 (vii) IMMEDIATE SOURCE: Synthetically Prepared  
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390 (viii) POSITION IN GENOME: None  
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392 (ix) FEATURE: None  
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394 (x) PUBLICATION INFORMATION: None  
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396 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:  
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398 GGAGAA 6  
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404 (i) SEQUENCE CHARACTERISTICS:  
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406 (A) LENGTH: 19  
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408 (B) TYPE: nucleic acid  
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410 (C) STRANDEDNESS: Single Stranded  
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412 (D) TOPOLOGY: Linear  
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414 (ii) MOLECULE TYPE: Not Applicable  
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416 (iii) HYPOTHETICAL: Not Applicable



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418 (iv) ANTI-SENSE: Not Applicable  
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420 (v) FRAGMENT TYPE: Not Applicable  
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422 (vi) ORIGINAL SOURCE: Synthetically Prepared  
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426 (viii) POSITION IN GENOME: None  
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428 (ix) FEATURE: None  
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432 (x) PUBLICATION INFORMATION: None  
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434 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:  
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436 CTCCTGAGGA GAAGTCTGC 19  
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439 (2) INFORMATION FOR SEQ ID NO:11:  
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441 (i) SEQUENCE CHARACTERISTICS:  
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443 (A) LENGTH: 19  
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445 (B) TYPE: nucleic acid  
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447 (C) STRANDEDNESS: Single Stranded  
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449 (D) TOPOLOGY: Linear  
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451 (ii) MOLECULE TYPE: Not Applicable  
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453 (iii) HYPOTHETICAL: Not Applicable  
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455 (iv) ANTI-SENSE: Not Applicable  
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457 (v) FRAGMENT TYPE: Not Applicable  
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459 (vi) ORIGINAL SOURCE: Synthetically Prepared  
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461 (vii) IMMEDIATE SOURCE: Synthetically Prepared  
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463 (viii) POSITION IN GENOME: None  
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465 (ix) FEATURE: None  
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467 (x) PUBLICATION INFORMATION: None  
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469 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:  
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471 CTCCTGTGGA GAAGTCTGC 19  
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475 (2) INFORMATION FOR SEQ ID NO:12:  
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477 (i) SEQUENCE CHARACTERISTICS:  
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479 (A) LENGTH: 23  
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481 (B) TYPE: nucleic acid  
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483 (C) STRANDEDNESS: Single Stranded  
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485 (D) TOPOLOGY: Linear  
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487 (ii) MOLECULE TYPE: Not Applicable  
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489 (iii) HYPOTHETICAL: Not Applicable  
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491 (iv) ANTI-SENSE: Not Applicable  
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493 (v) FRAGMENT TYPE: Not Applicable  
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495 (vi) ORIGINAL SOURCE: Synthetically Prepared  
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497 (vii) IMMEDIATE SOURCE: Synthetically Prepared  
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499 (viii) POSITION IN GENOME: None  
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501 (ix) FEATURE: None  
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503 (x) PUBLICATION INFORMATION: None  
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511 (2) INFORMATION FOR SEQ ID NO:13:  
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513 (i) SEQUENCE CHARACTERISTICS:  
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515 (A) LENGTH: 23  
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517 (B) TYPE: nucleic acid  
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519 (C) STRANDEDNESS: Single Stranded  
520

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521 (D) TOPOLOGY: Linear  
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523 (ii) MOLECULE TYPE: Not Applicable  
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525 (iii) HYPOTHETICAL: Not Applicable  
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527 (iv) ANTI-SENSE: Not Applicable  
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529 (v) FRAGMENT TYPE: Not Applicable  
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537 (ix) FEATURE: None  
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539 (x) PUBLICATION INFORMATION: None  
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549 (i) SEQUENCE CHARACTERISTICS:  
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551 (A) LENGTH: 14  
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553 (B) TYPE: nucleic acid  
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555 (C) STRANDEDNESS: Single Stranded  
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557 (D) TOPOLOGY: Linear  
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559 (ii) MOLECULE TYPE: Not Applicable  
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567 (vi) ORIGINAL SOURCE: Synthetically Prepared  
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569 (vii) IMMEDIATE SOURCE: Synthetically Prepared  
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571 (viii) POSITION IN GENOME: None  
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## Patent Application US/07/996,771

573 (ix) FEATURE: None  
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575 (x) PUBLICATION INFORMATION: None  
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577 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:  
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587 (A) LENGTH: 8  
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589 (B) TYPE: nucleic acid  
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591 (C) STRANDEDNESS: Single Stranded  
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593 (D) TOPOLOGY: Linear  
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595 (ii) MOLECULE TYPE: Not Applicable  
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607 (viii) POSITION IN GENOME: None  
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SEQUENCE VERIFICATION REPORT  
PATENT APPLICATION US/07/996,771

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LINE ERROR

ORIGINAL TEXT

51 Unknown or Misplaced Identifier  
58 Unknown or Misplaced Identifier

(C) CLASSIFICATION: Unknown  
(C) CLASSIFICATION: Unknown

PAGE: 1

SEQUENCE MISSING ITEM REPORT  
PATENT APPLICATION US/07/996,771

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S381

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SEQUENCE CORRECTION REPORT  
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S381

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CORRECTED TEXT